



OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/905,173

DATE: 05/20/2002
TIME: 14:07:46

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\05202002\I905173.raw

2 <110> APPLICANT: DIVERSA CORPORATION
3 SHORT, Jay M.
4 WARREN, Patrick V.
5 SWANSON, Ronald V.
6 MATHUR, Eric J.
8 <120> TITLE OF INVENTION: ENZYMES HAVING TRANSAMINASE AND AMINOTRANSFERASE ACTIVITY
AND METHODS OF
9 USE THEREOF
11 <130> FILE REFERENCE: DIVER1240-7
13 <140> CURRENT APPLICATION NUMBER: US 09/905,173
14 <141> CURRENT FILING DATE: 2001-07-12
16 <150> PRIOR APPLICATION NUMBER: US 09/412,184
17 <151> PRIOR FILING DATE: 1999-10-04
19 <150> PRIOR APPLICATION NUMBER: US 09/389,537
20 <151> PRIOR FILING DATE: 1999-09-02
22 <150> PRIOR APPLICATION NUMBER: US 08/646,590
23 <151> PRIOR FILING DATE: 1996-05-08
25 <150> PRIOR APPLICATION NUMBER: US 08/599,171
26 <151> PRIOR FILING DATE: 1996-02-09
28 <150> PRIOR APPLICATION NUMBER: US 09/481,733
29 <151> PRIOR FILING DATE: 2000-01-11
31 <150> PRIOR APPLICATION NUMBER: US 09/069,226
32 <151> PRIOR FILING DATE: 1998-04-27
34 <160> NUMBER OF SEQ ID NOS: 40
36 <170> SOFTWARE: PatentIn version 3.0
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62 <210> SEQ ID NO: 3
63 <211> LENGTH: 52
64 <212> TYPE: DNA

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240 cttggtatgg gcaatcctaa catgcctcca gcaaagcaca taatagataa actctgcgaa      180
242 gtggctcaaa agccgaacgt tcacggatat tctgcgtcaa ggggcatacc aagactgaga      240
244 aaggctatat gtaacttcta cgaagaaagg tacggagtga aactcgaccc tgagaggagg      300
246 gctatactaa caatcgggtgc aaaggaaggg tattctcatt tgatgcttgc gatgatattc      360
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250 attgcaggag gggaagttca ctcaataccc cttaacttct cggacgatca agatcatcag      480
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258 gatatagcct ttgacgggta caagcccccc tcaatactcg aaatagaagg tgctaaagac      720
260 gttgcgggtg agctctactc catgtcaaa ggccttttcaa tggcgggctg gagggtagcc      780
262 tttgtcgttg gaaacgaaat actcataaaa aaccttgcac acctcaaaag ctacttggat      840
264 tacggtatat ttactcccat acagggtggc tctattatcg cattagagag cccctacgaa      900
266 atcgtggaaa aaaccgcaaa ggtttaccaa aaaagaagag acgttctggt ggaagggtta      960
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270 cccgaatgga taaatatgaa ctctctggac ttttcttgt tctcctaaa agaggcgaag      1080
272 gttgcggtat ccccggtgt gggcttttgt cagtacggag aggggtacgt aaggtttgca      1140
274 cttgtagaaa atgaacacag gatcagacag gctataagg gataaaggaa agccttcaga      1200
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289 gtaatggaag ctctggaacg tgcggtgaag gaaaagacgt tcttctacac cctgctctg      180
291 ggactctggg aactcaggga aaggatatcg gagttttaca ggaaaaagta cagcgttgaa      240
293 gtttctccag agagagtcac cgtaactacc ggaacttcgg gagcgtttct cgtagcctac      300
295 gccgtaacac taaatgcggg agagaagata atcctcccag acccctctta cccctgttac      360
297 aaaaactttg cctacctctt agacgctcag ccggttttcg taaacgttga caaggaaacg      420

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299 aattacgaag taaggaaaga gatgatagaa gacattgatg cgaaagccct tcacatttcc 480
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303 tactgcgaag agaagggtat gtacttcata tccgacgaga ttaccacgg actcgtttac 600
305 gaaggtaggg agcacacagc acttgagttc tctgacaggg ctattgtcat aaacgggttt 660
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309 gtgagaaagg cggaaatagt aattcagaac gtatttatat ctgccccgac gctcagtcag 780
311 tacgccgccc ttgaggcttt tgattacgag tatttgagga aggtaagaaa aacctttgaa 840
313 gagaggagga acttccttta tggggaactg aaaaaactct tcaagataga cgcgaaacct 900
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317 gctttaaaac ttttaaggga ggcgaggggtg gcggtaacgc ccgggggtgga ctttgaaaaa 1020
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334 gtttacctgt gggacatata cggcaggaag tatatagatg ccatactctt cctctggtgc 180
336 aacgtccacg gacataacca ccctaaactg aacaacgcag ttatgaaaca gctctgtaag 240
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344 aaaaacgttt tcataacgct ttccgaagcc taccacgggg atactgtagg agcgggttagc 480
346 gtagggggta tagaactctt ccacggaact tataaagatc tccttttcaa gactataaaa 540
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352 attatggaag cgggaattca ggcagccgcg ggaatgctcc cttccctcc gggatttttg 720
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364 tttgaggaag aaagaacttt agagaagctc caaccaaaga taaagctttt aaaggaaagg 1080
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368 ggaatagagc tggtagagga caaagaaaag ggagaacctt tcccttacgg tgaaaggacg 1200
370 ggatttaagg tggcttacia gtgcagggaa aaaggggtgt ttttgagacc gctcggagac 1260
372 gttatggtat tgatgatgcc tcttgtaata gaggaagacg aaatgaacta cgttattgat 1320
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385 gtttacctgt acgatgagga aggaaaggag tatcttgact ttgtctccgg tataggcgctc 120
387 aactccctcg gtcacgctta cccaaaactc acagaagctc taaaagaaca ggttgagaaa 180
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:35; N Pos. 986
Seq#:36; Xaa Pos. 329

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 8